



CORRECTED SEQUENCE LISTING

<110> Masakazu Takeuchi
Yoshimi Takai

<120> Rabconnectin-3-binding protein

<130> 2003946-0055

<140> 10/645,335

<141> 2003-08-21

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<151> 2002-11-01

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gcg ccc aca cat tgc atc tca gcg gta ctt tta aca gat gat ggg gcc	96
Ala Pro Thr His Cys Ile Ser Ala Val Leu Leu Thr Asp Asp Gly Ala	
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acg atc gta aca gga tgt cac gac gga caa ata tgt ctc tgg gat ctt	144
Thr Ile Val Thr Gly Cys His Asp Gly Gln Ile Cys Leu Trp Asp Leu	
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tca gta gaa ctg caa att aat cct cga gca ctg ttg ttt ggt cat aca	192
Ser Val Glu Leu Gln Ile Asn Pro Arg Ala Leu Leu Phe Gly His Thr	
50 55 60	
gca tca atc act tgt ttg tct aaa gct tgt gct tcc agt gac aaa cag	240
Ala Ser Ile Thr Cys Leu Ser Lys Ala Cys Ala Ser Ser Asp Lys Gln	
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tat att gtg agt gca tct gaa agt gga gag atg tgc ctc tgg gat gtg	288
Tyr Ile Val Ser Ala Ser Glu Ser Gly Glu Met Cys Leu Trp Asp Val	
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agt gat ggc aga tgt att gaa ttt aca aaa tta gct tgc aca cat act	336
Ser Asp Gly Arg Cys Ile Glu Phe Thr Lys Leu Ala Cys Thr His Thr	

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Gly Ile Gln Phe Tyr Gln Phe Ser Val Gly Asn Gln Arg Glu Gly Arg			
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Leu Leu Cys His Gly His Tyr Pro Glu Ile Leu Val Val Asp Ala Thr			
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Ser Leu Glu Val Leu Tyr Ser Leu Val Ser Lys Ile Ser Pro Asp Trp			
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Ile Ser Ser Met Ser Ile Ile Arg Ser His Arg Thr Gln Glu Asp Thr			
165	170	175	
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Val Val Ala Leu Ser Val Thr Gly Ile Leu Lys Val Trp Ile Val Thr			
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tcg gaa ata agt gac atg cag gat act gag cca ata ttt gag gag gaa			624
Ser Glu Ile Ser Asp Met Gln Asp Thr Glu Pro Ile Phe Glu Glu Glu			
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210	215	220	
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Phe Thr Gln Arg Ser Leu Leu Val Val Cys Ser Lys Tyr Trp Arg Val			
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Gly Gln Thr Trp Thr Gly Gly Asp Phe Val Ser Ser Asp Lys Val Ile			
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Cys Leu Pro Ala Ser Asp Ser Phe Arg Ser Asp Val Gly Lys Ala Val			
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gaa aat tta att cct cct gta caa cat atc ctc ttg gat cga aaa gat			960
Glu Asn Leu Ile Pro Pro Val Gln His Ile Leu Leu Asp Arg Lys Asp			
305	310	315	320
aaa gag ttg cta att tgt cct cct gtt act cgg ttc ttc tat gga tgc			1008
Lys Glu Leu Leu Ile Cys Pro Pro Val Thr Arg Phe Phe Tyr Gly Cys			
325	330	335	
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Arg Glu Tyr Phe His Lys Leu Leu Ile Gln Gly Asp Ser Ser Gly Arg			
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Glu Gly Leu Ala Met Thr Thr Ser Ile Ser Leu Gln Glu Ala Phe Asp	
370 375 380	
aaa ctg aat cct tgt cct gct gga att ata gat cag ctg agt gtg att	1200
Lys Leu Asn Pro Cys Pro Ala Gly Ile Ile Asp Gln Leu Ser Val Ile	
385 390 395 400	
ccc aat agt aat gaa cct ctt aaa gta act gca agt gtg tac ata cca	1248
Pro Asn Ser Asn Glu Pro Leu Lys Val Thr Ala Ser Val Tyr Ile Pro	
405 410 415	
gca cat gga cga ctt gtt tgt ggt cgt gaa gat gga agc ata gtt att	1296
Ala His Gly Arg Leu Val Cys Gly Arg Glu Asp Gly Ser Ile Val Ile	
420 425 430	
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Val Pro Ala Thr Gln Thr Ala Ile Val Gln Leu Leu Gln Gly Glu His	
435 440 445	
atg ctc aga aga ggt tgg cca cct cac aga aca ctc cgt ggt cat cgg	1392
Met Leu Arg Arg Gly Trp Pro Pro His Arg Thr Leu Arg Gly His Arg	
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Asn Lys Val Thr Cys Leu Leu Tyr Pro His Gln Val Ser Ala Arg Tyr	
465 470 475 480	
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Asp Gln Arg Tyr Leu Ile Ser Gly Gly Val Asp Phe Ser Val Ile Ile	
485 490 495	
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Trp Asp Ile Phe Ser Gly Glu Met Lys His Ile Phe Cys Val His Gly	
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Gly Glu Ile Thr Gln Leu Leu Val Pro Pro Glu Asn Cys Ser Ala Arg	
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Val Gln His Cys Ile Cys Ser Val Ala Ser Asp His Ser Val Gly Leu	
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Phe Pro Ile Gln Val Ile Lys Trp Arg Pro Ser Asp Asp Tyr Leu Val	
565 570 575	
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Val Gly Cys Ser Asp Gly Ser Val Tyr Val Trp Gln Met Asp Thr Gly	
580 585 590	
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Ala Leu Asp Arg Cys Val Met Gly Ile Thr Ala Val Glu Ile Leu Asn	
595 600 605	

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Ala Cys Asp Glu Ala Val Pro Ala Ala Val Asp Ser Leu Ser His Pro	
610 615 620	
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Ala Val Asn Leu Lys Gln Ala Met Thr Arg Arg Ser Leu Ala Ala Leu	
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Lys Asn Met Ala His His Lys Leu Gln Thr Leu Ala Thr Asn Leu Leu	
645 650 655	
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Ala Ser Glu Ala Ser Asp Lys Gly Asn Leu Pro Lys Tyr Ser His Asn	
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Ser Leu Met Val Gln Ala Ile Lys Thr Asn Leu Thr Asp Pro Asp Ile	
675 680 685	
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His Val Leu Phe Phe Asp Val Glu Ala Leu Ile Ile Gln Leu Leu Thr	
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caa aaa gca tct ggc agt tca gac aaa ggg ggc tct ttt tta act gga	2208
Gln Lys Ala Ser Gly Ser Ser Asp Lys Gly Gly Ser Phe Leu Thr Gly	
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Asn Ile Lys Glu His Leu Leu Asp Asp Glu Glu Glu Asp Glu Glu Ile	
755 760 765	
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Met Arg Gln Arg Arg Glu Glu Ser Asp Pro Glu Tyr Arg Ser Ser Lys	
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Ser Lys Pro Leu Thr Leu Leu Glu Tyr Asn Leu Thr Met Asp Thr Ala	
785 790 795 800	
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Lys Leu Phe Met Ser Cys Leu His Ala Trp Gly Leu Asn Glu Val Leu	
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Asp Glu Val Cys Leu Asp Arg Leu Gly Met Leu Lys Pro His Cys Thr	
820 825 830	
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Val Ser Phe Gly Leu Leu Ser Arg Gly Gly His Met Ser Leu Met Leu	
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Pro Gly Tyr Asn Gln Pro Ala Cys Lys Leu Ser His Gly Lys Thr Glu	

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Val Gly Arg Lys Leu Pro Ala Ser Glu Gly Val Gly Lys Gly Thr Tyr			
865	870	875	880
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Gly Val Ser Arg Ala Val Thr Thr Gln His Leu Leu Ser Ile Ile Ser			
885	890	895	
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Leu Ala Asn Thr Leu Met Ser Met Thr Asn Ala Thr Phe Ile Gly Asp			
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cat atg aag aag ggt cct acc agg cca cct aga cca agc acc cca gac			2784
His Met Lys Lys Gly Pro Thr Arg Pro Pro Arg Pro Ser Thr Pro Asp			
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ctt tct aag gca agg ggt tcc cct cca act tcc agt aat att gtg caa			2832
Leu Ser Lys Ala Arg Gly Ser Pro Pro Thr Ser Ser Asn Ile Val Gln			
930	935	940	
gga cag att aaa caa gtt gct gca cct gtc gtt tcc gct cgg tct gat			2880
Gly Gln Ile Lys Gln Val Ala Ala Pro Val Val Ser Ala Arg Ser Asp			
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Val Met Leu Pro Asp Leu Leu Gly Leu Asp Lys Phe Arg Pro Pro Leu			
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ctg gag atg ctg gcc cga aga tgg caa gat cga tgc ttg gag gtg			3069
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Arg Glu Ala Ala Gln Ala Leu Leu Leu Ala Glu Leu Arg Arg Ile			
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Leu Pro Gln Tyr Ile Asp His Val Ile Ser Pro Gly Val Thr Ser			
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Glu Ala Ala Gln Thr Ile Thr Thr Ala Pro Asp Ala Ser Gly Pro			
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Pro Lys Leu Leu Thr Arg Pro Arg Ser Ser Ser Gln Ile Pro Glu	
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Leu Ile Gly Arg Gly Phe Thr Val Trp Glu Pro Tyr Met Asp Val	
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Ser Ala Val Leu Met Gly Leu Leu Glu Leu Cys Ala Asp Ala Glu	
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Thr Ala Arg Pro Pro Ala Phe Ile Thr Thr Ile Ala Lys Glu Val	
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Met His Thr Thr Thr Leu Ala Arg Ala Lys Gly Glu Ile Leu Arg	
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Val Ile Glu Ile Leu Ile Glu Lys Met Pro Thr Asp Val Val Asp	
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Ala Val Ala Phe Ala Pro Asp Gly Arg Tyr Leu Ala Thr Tyr Ser	
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Asn Thr Asp Ser His Ile Ser Phe Trp Gln Met Asn Thr Ser Leu	
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cgc aac gtc atc ctc atg gcc cat gac ggg aag gag cac cgc ttc	4464
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Met Val	
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Thr Ile Val Thr Gly Cys His Asp Gly Gln Ile Cys Leu Trp Asp Leu	
35 40 45	
Ser Val Glu Leu Gln Ile Asn Pro Arg Ala Leu Leu Phe Gly His Thr	

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Ala Ser Ile Thr Cys	Leu Ser Lys Ala Cys	Ala Ser Ser Asp Lys Gln
65	70	75
Tyr Ile Val Ser Ala	Ser Glu Ser Gly Glu Met	Cys Leu Trp Asp Val
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Ser Asp Gly Arg Cys	Ile Glu Phe Thr Lys	Leu Ala Cys Thr His Thr
100	105	110
Gly Ile Gln Phe Tyr	Gln Phe Ser Val Gly	Asn Gln Arg Glu Gly Arg
115	120	125
Leu Leu Cys His Gly	His Tyr Pro Glu Ile	Leu Val Val Asp Ala Thr
130	135	140
Ser Leu Glu Val Leu	Tyr Ser Leu Val Ser	Lys Ile Ser Pro Asp Trp
145	150	155
Ile Ser Ser Met Ser	Ile Ile Arg Ser His	Arg Thr Gln Glu Asp Thr
165	170	175
Val Val Ala Leu Ser	Val Thr Gly Ile Leu	Lys Val Trp Ile Val Thr
180	185	190
Ser Glu Ile Ser Asp	Met Gln Asp Thr Glu	Pro Ile Phe Glu Glu Glu
195	200	205
Ser Lys Pro Ile Tyr	Cys Gln Asn Cys Gln	Ser Ile Ser Phe Cys Ala
210	215	220
Phe Thr Gln Arg Ser	Leu Leu Val Val Cys	Ser Lys Tyr Trp Arg Val
225	230	235
Phe Asp Ala Gly Asp	Tyr Ser Leu Leu Cys	Ser Gly Pro Ser Glu Asn
245	250	255
Gly Gln Thr Trp Thr	Gly Gly Asp Phe Val	Ser Ser Asp Lys Val Ile
260	265	270
Ile Trp Thr Glu Asn	Gly Gln Ser Tyr Ile	Tyr Lys Leu Pro Ala Ser
275	280	285
Cys Leu Pro Ala Ser	Asp Ser Phe Arg Ser	Asp Val Gly Lys Ala Val
290	295	300
Glu Asn Leu Ile Pro	Pro Val Gln His Ile	Leu Leu Asp Arg Lys Asp
305	310	315
Lys Glu Leu Leu Ile	Cys Pro Pro Val Thr	Arg Phe Phe Tyr Gly Cys
325	330	335
Arg Glu Tyr Phe His	Lys Leu Leu Ile Gln	Gly Asp Ser Ser Gly Arg
340	345	350
Leu Asn Ile Trp Asn	Ile Ser Asp Thr Ala	Asp Lys Gln Gly Ser Glu
355	360	365
Glu Gly Leu Ala Met	Thr Thr Ser Ile Ser	Leu Gln Glu Ala Phe Asp
370	375	380
Lys Leu Asn Pro Cys	Pro Ala Gly Ile Ile	Asp Gln Leu Ser Val Ile
385	390	395
Pro Asn Ser Asn Glu	Pro Leu Lys Val Thr	Ala Ser Val Tyr Ile Pro
405	410	415
Ala His Gly Arg Leu	Val Cys Gly Arg Glu	Asp Gly Ser Ile Val Ile
420	425	430

Val Pro Ala Thr Gln Thr Ala Ile Val Gln Leu Leu Gln Gly Glu His
 435 440 445
 Met Leu Arg Arg Gly Trp Pro Pro His Arg Thr Leu Arg Gly His Arg
 450 455 460
 Asn Lys Val Thr Cys Leu Leu Tyr Pro His Gln Val Ser Ala Arg Tyr
 465 470 475 480
 Asp Gln Arg Tyr Leu Ile Ser Gly Gly Val Asp Phe Ser Val Ile Ile
 485 490 495
 Trp Asp Ile Phe Ser Gly Glu Met Lys His Ile Phe Cys Val His Gly
 500 505 510
 Gly Glu Ile Thr Gln Leu Leu Val Pro Pro Glu Asn Cys Ser Ala Arg
 515 520 525
 Val Gln His Cys Ile Cys Ser Val Ala Ser Asp His Ser Val Gly Leu
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 Leu Ser Leu Arg Glu Lys Lys Cys Ile Met Leu Ala Ser Arg His Leu
 545 550 555 560
 Phe Pro Ile Gln Val Ile Lys Trp Arg Pro Ser Asp Asp Tyr Leu Val
 565 570 575
 Val Gly Cys Ser Asp Gly Ser Val Tyr Val Trp Gln Met Asp Thr Gly
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 Ala Leu Asp Arg Cys Val Met Gly Ile Thr Ala Val Glu Ile Leu Asn
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 Ala Cys Asp Glu Ala Val Pro Ala Ala Val Asp Ser Leu Ser His Pro
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 Ala Val Asn Leu Lys Gln Ala Met Thr Arg Arg Ser Leu Ala Ala Leu
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 Lys Asn Met Ala His His Lys Leu Gln Thr Leu Ala Thr Asn Leu Leu
 645 650 655
 Ala Ser Glu Ala Ser Asp Lys Gly Asn Leu Pro Lys Tyr Ser His Asn
 660 665 670
 Ser Leu Met Val Gln Ala Ile Lys Thr Asn Leu Thr Asp Pro Asp Ile
 675 680 685
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 Glu Glu Ala Ser Arg Pro Asn Thr Ala Leu Ile Ser Pro Glu Asn Leu
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 Lys Arg Ala Ala Val Leu Phe Gln Gln Val Lys Glu Thr Ile Lys Glu
 740 745 750
 Asn Ile Lys Glu His Leu Leu Asp Asp Glu Glu Glu Asp Glu Glu Ile
 755 760 765
 Met Arg Gln Arg Arg Glu Glu Ser Asp Pro Glu Tyr Arg Ser Ser Lys
 770 775 780
 Ser Lys Pro Leu Thr Leu Leu Glu Tyr Asn Leu Thr Met Asp Thr Ala
 785 790 795 800
 Lys Leu Phe Met Ser Cys Leu His Ala Trp Gly Leu Asn Glu Val Leu

	805		810		815
Asp Glu Val Cys Leu Asp Arg Leu Gly Met Leu Lys Pro His Cys Thr					
	820		825		830
Val Ser Phe Gly Leu Leu Ser Arg Gly Gly His Met Ser Leu Met Leu					
	835		840		845
Pro Gly Tyr Asn Gln Pro Ala Cys Lys Leu Ser His Gly Lys Thr Glu					
	850		855		860
Val Gly Arg Lys Leu Pro Ala Ser Glu Gly Val Gly Lys Gly Thr Tyr					
865		870		875	880
Gly Val Ser Arg Ala Val Thr Thr Gln His Leu Leu Ser Ile Ile Ser					
	885		890		895
Leu Ala Asn Thr Leu Met Ser Met Thr Asn Ala Thr Phe Ile Gly Asp					
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His Met Lys Lys Gly Pro Thr Arg Pro Pro Arg Pro Ser Thr Pro Asp					
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Leu Ser Lys Ala Arg Gly Ser Pro Pro Thr Ser Ser Asn Ile Val Gln					
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Gly Gln Ile Lys Gln Val Ala Ala Pro Val Val Ser Ala Arg Ser Asp					
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Ala Asp His Ser Gly Ser Asp Pro Pro Ser Ala Pro Ala Leu His Thr					
	965		970		975
Cys Phe Leu Val Asn Glu Gly Trp Ser Gln Leu Ala Ala Met His Cys					
	980		985		990
Val Met Leu Pro Asp Leu Leu Gly Leu Asp Lys Phe Arg Pro Pro Leu					
	995		1000		1005
Leu Glu Met Leu Ala Arg Arg Trp Gln Asp Arg Cys Leu Glu Val Arg					
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Glu Ala Ala Gln Ala Leu Leu Leu Ala Glu Leu Arg Arg Ile Glu Gln					
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Ala Gly Arg Lys Glu Ala Ile Asp Ala Trp Ala Pro Tyr Leu Pro Gln					
	1045		1050		1055
Tyr Ile Asp His Val Ile Ser Pro Gly Val Thr Ser Glu Ala Ala Gln					
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Thr Ile Thr Thr Ala Pro Asp Ala Ser Gly Pro Glu Ala Lys Val Gln					
	1075		1080		1085
Glu Glu Glu His Asp Leu Val Asp Asp Asp Ile Thr Thr Gly Cys Leu					
	1090		1095		1100
Ser Ser Val Pro Gln Met Lys Lys Ile Ser Thr Ser Tyr Glu Glu Arg					
1105		1110		1115	1120
Arg Lys Gln Ala Thr Ala Ile Val Leu Leu Gly Val Ile Gly Ala Glu					
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Phe Gly Ala Glu Ile Glu Pro Pro Lys Leu Leu Thr Arg Pro Arg Ser					
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Ser Ser Gln Ile Pro Glu Gly Phe Gly Leu Thr Ser Gly Gly Ser Asn					
	1155		1160		1165
Tyr Ser Leu Ala Arg His Thr Cys Lys Ala Leu Thr Phe Leu Leu Leu					
	1170		1175		1180

Gln Pro Pro Ser Pro Lys Leu Pro Pro His Ser Thr Ile Arg Arg Thr
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 Ala Ile Asp Leu Ile Gly Arg Gly Phe Thr Val Trp Glu Pro Tyr Met
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 Asp Val Ser Ala Val Leu Met Gly Leu Leu Glu Leu Cys Ala Asp Ala
 1220 1225 1230
 Glu Lys Gln Leu Ala Asn Ile Thr Met Gly Leu Pro Leu Ser Pro Ala
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 Ala Asp Ser Ala Arg Ser Ala Arg His Ala Leu Ser Leu Ile Ala Thr
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 Ala Arg Pro Pro Ala Phe Ile Thr Thr Ile Ala Lys Glu Val His Arg
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 His Thr Ala Leu Ala Ala Asn Thr Gln Ser Gln Gln Asn Met His Thr
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 Thr Thr Leu Ala Arg Ala Lys Gly Glu Ile Leu Arg Val Ile Glu Ile
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 Leu Ile Glu Lys Met Pro Thr Asp Val Val Asp Leu Leu Val Glu Val
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 Met Asp Ile Ile Met Tyr Cys Leu Glu Gly Ser Leu Val Lys Lys Lys
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 Gly Leu Gln Glu Cys Phe Pro Ala Ile Cys Arg Phe Tyr Met Val Ser
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 Tyr Tyr Glu Arg Asn His Arg Ile Ala Val Gly Ala Arg His Gly Ser
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 Val Ala Leu Tyr Asp Ile Arg Thr Gly Lys Cys Gln Thr Ile His Gly
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 His Lys Gly Pro Ile Thr Ala Val Ala Phe Ala Pro Asp Gly Arg Tyr
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